Please amend the specification as follows:

Please replace the paragraph beginning at page 20, line 8 with the following amended paragraph:

--- Herpes virus entry mediator (HVEM) is a TNF family receptor related to the LT.beta.R and binds tightly to the ligand LIGHT tightly and weakly to the ligand lymphotoxin-a (LTa) (Mauri et al, 1998). Human HVEM was prepared as a Ig fusion protein by PCR amplification of the extracellular domain and fused to the human IgG1 CH2 and CH3 region as described for LT.beta.R-Ig (Crowe et al, 1994). The construct was inserted into a vector called CH269 (Chicheportiche et al, 1997) for transient expression in the human embryonic kidney cell line 293 with high copy vector expression using the EBNA system (293-E cells). Supernatents were collected and HVEM-Ig purified using ProteinA affinity chromatography and low pH elution. Recombinant LT.alpha. was prepared from insect cells as described (Browning et al, 1996a). Recombinant soluble human LIGHT was prepared by PCR amplification of the entire cDNA using RNA from activated II23 cells yielding the coding region of the sequence described by Mauri et al, 1998. The receptor binding domain of LIGHT was amplified by PCR and fused onto the alpha mating factor leader sequence and expressed essentially as described for other related proteins (Browning et al, 1996). A FLAG tag and (G.sub.4S).sub.3 spacer amino acid sequences were inserted between the leader and the receptor binding domain such that the secreted LIGHT would possess a N-terminal FLAG sequence. The construct encoded the following molecule: "MRFPSIFTAVLFAASSALAAPVNTT TEDETAQIPAEAVIGYSDLEGDFDVAVLPFSNSTNNGLLFINTTIASIAAKEEGVSLEK R..EADYKDDDDNGGGSGGGSGGGSKELNPAAHLTGANSSLTGSGGPLLWETQLGLA FLRGLSYHDGALVVTKAGYYYIYSKVQLGGVGCPLGLASTITHGLYKRTPRYPEELE LLVSQQSPCGRATSSSRVWWDSSFLGGVVHLEAGEEVVVRVLDERLVRLRDGTRSY FGAFMV" (SEQ ID NO: 1) where the two dots indicates the expected N-terminus of the mature protein which could be further processed by removal of the next two amino acids (EA). The protein was purified from the supernatant by affinity chormatography over an anti-FLAG mAb column and elution with either low pH or calcium chelation. Full length LIGHT was also inserted into a vector CH269 for expression on the surface of 293-E cells as

described (Chicheportiche et al, 1997). FACS binding methods for the detection of receptor-Ig to cell surfaces and BIAcore methods for measuring the binding of soluble ligands to immobilized receptor-Ig have been described (Mackay et al, 1997). The BIAcore technology yields real time measurements of protein bound to the chips (i.e. the receptor). ---